

**In the Specification:**

Please amend the specification as shown:

Please delete paragraph [0001] and replace it with the following paragraph:

**[0001]** This application claims priority to U.S. Provisional Application No. as 60/339,466, filed February 27, 2003, U.S. Application Serial No. 09/989,497, filed November 21, 2001, which is a continuation-in-part of U.S. Application Serial No. 09/984,292, filed October 29, 2001, which claims priority to U.S. Provisional Application No. 60/243,770, filed on October 30, 2000, and all of these applications are incorporated herein by reference in their entirety.

Please delete paragraph [0090] and replace it with the following paragraph:

**[0090]** In this example, the details of how the Ga15 chimeras were made are presented. Ga<sub>15</sub> chimeras were generated by PCR with mutagenic 3' primers. The sequence of the parental Ga15clone (SEQ ID NO:1; amino acid sequence is SEQ ID NO:2) corresponds to databank sequences (e.g., accession B0005439) except for a silent single nucleotide polymorphism. The last six codons of GOC15 and the sequences they were replaced with are shown below (SEQ ID NOS:3-12). The Ga15 chimeras were generated with 5' Ascl sites (GGCGCGCCGCC (SEQ ID NO: 13) joined to the start ATG) and 3' NotI sites (GCGGCCGC joined to the stop TGA) and cloned as Ascl-NotI fragments in the Ascl-NotI polylinker sites of the pEAK10 expression vector (Edge Biosystems).

**Ga15 nucleotide sequence (SEQ ID NO:1)**

atggccccggtccctgacttggggctgctgtccctggtgcctgacagaggaggagaagactgccgccagaa  
tcgaccaggagatcaacaggattttgttggaacagaaaaacaagagcgcgaggaattgaaactcctgct  
gttggggcctggtgagagcgggaagagtacgttcatcaagcagatgcgcattcattcacgggtgtgggctac  
tcggaggaggaccgcagagccttcgggctgctcatctaccagaacatcttcgtctccatgcaggccatga  
tagatgcgatggaccggctgcagatcccccttcagcaggcctgacagcaagcagcagccagcctagtgat  
gaccaggagaccctataaagtgagcacattcgagaagccatatgcagtggccatgcagtaacctgtggcgg  
gacgcgggcatccgtgcatgctacgagcgaaggcgtgaattccaccttctggactccgcggtgtattacc  
tgtcacacctggagcgcataatcagaggacagctacatccccactgcgcaagacgtgctgcgcagtcgcat

gcccaccacagggcatcaatgagtactgcttctccgtgaagaaaaccaaactgcgcacgtggatgttggt  
ggccagaggtcagagcgtaggaaatggattcactgttttgagaacgtgattgccctcatctacctggcct  
ccctgagcaggtatgaccagtgcctagaggagaacgatcaggagaaccgcacgtggaggagagtctcgctct  
gttcagcacgatacctagagctgccctgggtcaagagcacctcgggtcatcctcttctcctcaacaagacggac  
atcctggaagataagattcacacctccacctggccacatacttccccagcttccagggacccccggcgag  
acgcagagggccgccaagagcttcatcttggacatgtatgcgcgcgtgtacgcgagctgcgcagagcccca  
ggacgggtggcaggaaggctcccgcgcgcgcgcgttcttcgcacacttcacctgtgccacggacacgcaa  
agcgtccgcagcgtgttcaaggacgtgcgggactcgggtgctggcccggtacctggacgagatcaacctgc  
tgtga

### **Ga<sub>15</sub>. amino acid sequence (SEQ ID NO:2)**

MARSLTWGCCPWCLTEEEKTAARIDQEINRILLEQKKQEREELKLLLLGPGESGKSTFIKQMRIIHGV  
GYSEEDRRAFRLLIYQNI FVSMQAMIDAMDRLQIPFSRPDSKQHASLVMTQDPYKVSTFEKPYAVAMQ  
YLWRDAGIRACYERRREFHLLDSAVYYLSHLERISEDSYIPTAQDVLRSRMPPTGINEYCFVSKTKL  
RIVDVGGQSRERRKWIHCFENVIALIYLASLSEYDQCLEENDQENRMEESLALFSTILELPWFKSTSV  
ILFLNKTDILEDKIHTSHLATYFPSFQGPRRDAAEAAKSFILDMYARVYASCAEPQDGGGRKGSRRARFF  
AHFTCATDTQSVRSVFKDVRDSVLARYLDEINLL

### **G<sub>a</sub> tails (SEQ IDS NOS: 3-12)**

GAGATCAACCTGCTGTGA	G <sub>a15</sub>	(SEQ ID NO: 3)
GACTGTGGCCTCTTCTGA	G <sub>ai1</sub>	(SEQ ID NO: 4)
GAGTACAATCTGGTCTGA	G <sub>aq</sub>	(SEQ ID NO: 5)
CAGTATGAGCTCTTGTGA	G <sub>as</sub>	(SEQ ID NO: 6)
GAGTGCGGCCTCTACTGA	G <sub>ai3</sub>	(SEQ ID NO: 7)
GGATGCGGACTCTACTGA	G <sub>ao</sub>	(SEQ ID NO: 8)
TACATCGGCCTCTGCTGA	G <sub>az</sub>	(SEQ ID NO: 9)
GACATCATGCTCCAATGA	G <sub>ai2</sub>	(SEQ ID NO: 10)
CAACTAATGCTCCAATGA	G <sub>a13</sub>	(SEQ ID NO: 11)
CACCAGGTTGAACTCTGA	G <sub>a14</sub>	(SEQ ID NO: 12)